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Search Results - <table border="1" style="display: inline-table; vertical-align: middle; border-collapse: collapse;"> <tr> <td style="padding: 2px 10px;">Terms</td> <td style="padding: 2px 10px;">Documents</td> </tr> <tr> <td style="padding: 2px 10px;">L3 AND L4</td> <td style="padding: 2px 10px;">1</td> </tr> </table>		Terms	Documents	L3 AND L4	1
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Search History

DATE: Friday, April 18, 2003 [Printable Copy](#) [Create Case](#)

Set Name	Query	Hit Count	Set Name
side by side			result set
DB=USPT,PGPB,JPAB,EPAB,DWPI; PLUR=YES; OP=ADJ			
<u>L6</u>	L3 AND L4	1	<u>L6</u>
<u>L5</u>	deb chatterjee.in.	0	<u>L5</u>
<u>L4</u>	chatterjee.in.	1438	<u>L4</u>
<u>L3</u>	maritima.clm.	40	<u>L3</u>
<u>L2</u>	maritima.clm. and chatterjee.in.	1	<u>L2</u>
<u>L1</u>	maritima and chatterjee.in.	15	<u>L1</u>

END OF SEARCH HISTORY

=> d 112 ibib ab 15-21

L12 ANSWER 15 OF 32 CAPLUS COPYRIGHT 2003 ACS
ACCESSION NUMBER: 1999:761460 CAPLUS
DOCUMENT NUMBER: 132:9599
TITLE: Detection of nucleic acids by multiple sequential
invasive cleavages
INVENTOR(S): Hall, Jeff G.; Lyamichev, Victor I.; Mast, Andrea L.;
Brow, Mary Ann D.
PATENT ASSIGNEE(S): Third Wave Technologies, Inc., USA
SOURCE: U.S., 306 pp., Cont.-in-part of U.S. Ser. No. 759,038.
DOCUMENT TYPE: CODEN: USXXAM
LANGUAGE: Patent
FAMILY ACC. NUM. COUNT: 12
PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 5994069	A	19991130	US 1997-823516	19970324
US 5846717	A	19981208	US 1996-599491	19960124
US 6001567	A	19991214	US 1996-682853	19960712
US 5985557	A	19991116	US 1996-756386	19961129
US 6090606	A	20000718	US 1996-758314	19961202
US 6090543	A	20000718	US 1996-759038	19961202
WO 9727214	A1	19970731	WO 1997-US1072	19970122
W: AU, CA, JP, US				
RW: AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE				
WO 9842873	A1	19981001	WO 1998-US5809	19980324
W: AU, CA, JP, US				
RW: AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE				
AU 9868690	A1	19981020	AU 1998-68690	19980324
AU 738849	B2	20010927		
EP 994964	A1	20000426	EP 1998-914299	19980324
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, FI				
JP 2001518805	T2	20011016	JP 1998-545934	19980324
US 6458535	B1	20021001	US 1999-350597	19990709
US 2002187486	A1	20021212	US 2001-33297	20011102
PRIORITY APPLN. INFO.:				
US 1996-599491 A2 19960124				
US 1996-682853 A2 19960712				
US 1996-756386 A2 19961129				
US 1996-758314 A2 19961202				
US 1996-759038 A2 19961202				
WO 1997-US1072 A2 19970122				
US 1996-756038 B2 19961126				
US 1996-756376 A2 19961202				
US 1997-823516 A2 19970324				
WO 1998-US5809 W 19980324				
US 1999-350597 A1 19990709				

AB The present invention relates to means for the detection and characterization of nucleic acid sequences, as well as variations in nucleic acid sequences, by an Invader.RTM. oligonucleotide-directed cleavage detection assay. The present invention also relates to methods for forming a nucleic acid cleavage structure on a target sequence and cleaving the nucleic acid cleavage structure in a site-specific manner. The structure-specific nuclease activity of a variety of enzymes is used to cleave the target-dependent cleavage structure, thereby indicating the presence of specific nucleic acid sequences or specific variations thereof. Derivs. of thermostable DNA polymerases and their mutants that retain their 5'-nuclease activity but lack polymerase activity are described for use in the nucleic acid detection system. The nuclease activity cleaves the single-stranded moiety of a Y-shaped structure and so is of use in selected cleavage of reporter

sequences in a hybridization assay that includes 5'-nuclease-dependent cleavage and amplification steps. The present invention further relates to methods and devices for the sepn. of nucleic acid mols. based on charge. The present invention also provides methods for the detection of non-target cleavage products via the formation of a complete and activated protein binding region. The invention further provides sensitive and specific methods for the detection of human cytomegalovirus nucleic acid in a sample.

REFERENCE COUNT: 14 THERE ARE 14 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L12 ANSWER 16 OF 32 CAPLUS COPYRIGHT 2003 ACS
ACCESSION NUMBER: 1999:64619 CAPLUS
DOCUMENT NUMBER: 130:121430
TITLE: **Mutant chimeric Thermus/Tma DNA polymerases** with improved properties for nucleic acid sequencing
INVENTOR(S): Gelfand, David Harrow; Reichert, Fred Lawrence
PATENT ASSIGNEE(S): F. Hoffmann-La Roche Ag, Switz.
SOURCE: Eur. Pat. Appl., 47 pp.
DOCUMENT TYPE: Patent
LANGUAGE: English
FAMILY ACC. NUM. COUNT: 1
PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
EP 892058	A2	19990120	EP 1998-112327	19980703
EP 892058	A3	20010926		
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO				
US 6228628	B1	20010508	US 1998-105697	19980626
CA 2240570	AA	19990109	CA 1998-2240570	19980707
JP 11089588	A2	19990406	JP 1998-208533	19980709
US 1997-52065P P 19970709				

PRIORITY APPLN. INFO.: AB The invention provides **mutant**, chimeric thermostable **DNA polymerase** enzymes consisting of an N-terminal region derived from the 5'-nuclease domain of a **Thermus** species **DNA polymerase** and a C-terminal region derived from the 3' to 5' exonuclease and polymerase domains of **Tma DNA polymerase**. These **mutant** chimeric thermostable **DNA polymerase** enzymes have improved properties in nucleic acid sequencing reactions. Also provided are nucleic acids encoding said **mutant** chimeric thermostable **DNA polymerase** enzymes, vectors comprising said nucleic acids and host cells transformed with said vectors. Also provided are compns. comprising said mutated, chimeric thermostable **DNA polymerase** enzymes and non-ionic polymeric detergent(s). Furthermore methods for producing the said enzymes and methods and kits for using the said enzymes are provided.

L12 ANSWER 17 OF 32 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC. DUPLICATE
1

ACCESSION NUMBER: 1999:236361 BIOSIS
DOCUMENT NUMBER: PREV199900236361
TITLE: New reagents for directed modification of biopolymers:
Photoaffinity modification of **Tte DNA polymerase**.
AUTHOR(S): Kolpashchikov, D. M.; Zakharenko, A. L.; Dezhurov, S. V.; Rechkunova, N. I.; Khodyreva, S. N.; Degtyarev, S. Kh.; Litvak, V. V.; Lavrik, O. I. (1)
CORPORATE SOURCE: (1) Siberian Division, Novosibirsk Institute of Bioorganic Chemistry, Russian Academy of Sciences, prosp. Akademika Lavrent'eva 8, Novosibirsk, 630090 Russia

SOURCE: Bioorganicheskaya Khimiya, (Feb., 1999) Vol. 25, No. 2, pp. 129-136.
 DOCUMENT TYPE: Article
 LANGUAGE: Russian
 SUMMARY LANGUAGE: English; Russian
 AB Arylazides N-(4-azido-2,5-difluoro-3-chloropyridinyl-6)-beta-alanine (Ia) and N-(4-azido-2,5-difluoro-3-chloropyridinyl-6)-glycine (Ib) were synthesized and covalently attached to 5-(3-aminopropenyl-1)-dUTP through the amino group to give 5'-triphosphate (IIa) and 5'-triphosphate (IIb). The resulting azides were subjected to photolysis in aqueous solution. The spectral and photochemical characteristics of azides (I) and (II) imply that their use for the modification of biopolymers holds promise. Compounds (IIa, b) effectively substituted dTTP in DNA polymerization catalyzed by thermostable DNA polymerase from *Thermus thermophilus* B-35 (Tte DNA polymerase). Photoaffinity modification of Tte DNA polymerase was carried out by dTTP analogues (IIa, b) and by earlier obtained 5-(N-(5-azido-2-nitrobenzoyl)-trans-3-aminopropenyl-1) deoxyuridine 5'-triphosphate (III) and 5-(N-(4-azido-2,3,5,6-tetrafluorobenzoyl)-trans-3-aminopropenyl-1) deoxyuridine 5'-triphosphate (IV) using two variants of labeling. All four dTTP analogues were shown to modify Tte DNA polymerase.

L12 ANSWER 18 OF 32 CAPLUS COPYRIGHT 2003 ACS
 ACCESSION NUMBER: 1998:388639 CAPLUS
 DOCUMENT NUMBER: 129:64906
 TITLE: Invasive cleavage of nucleic acids for detecting and characterizing target nucleic acids and microbial nucleases for the methods
 INVENTOR(S): Kaiser, Michael W.; Lyamichev, Victor I.; Lyamicheva, Natasha
 PATENT ASSIGNEE(S): Third Wave Technologies, Inc., USA; Kaiser, Michael W.; Lyamichev, Victor I.; Lyamicheva, Natasha
 SOURCE: PCT Int. Appl., 472 pp.
 DOCUMENT TYPE: Patent
 LANGUAGE: English
 FAMILY ACC. NUM. COUNT: 12
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9823774	A1	19980604	WO 1997-US21783	19971126
W: AU, CA, JP, US, US				
RW: AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE				
US 5843669	A	19981201	US 1996-757653	19961129
US 6090606	A	20000718	US 1996-758314	19961202
AU 9855898	A1	19980622	AU 1998-55898	19971126
AU 737449	B2	20010823		
EP 966542	A1	19991229	EP 1997-952237	19971126
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, FI				
JP 2001526526	T2	20011218	JP 1998-524043	19971126
PRIORITY APPLN. INFO.:			US 1996-757653	A2 19961129
			US 1996-758314	A2 19961202
			US 1996-599491	A2 19960124
			US 1996-682853	A2 19960712
			US 1996-756386	A2 19961129
			US 1996-756376	A2 19961202
			WO 1997-US21783	W 19971126

AB The present invention relates to means for the detection and characterization of nucleic acid sequences, as well as variations in nucleic acid sequences. The present invention also relates to improved

cleavage means for the detection and characterization of nucleic acid sequences. Structure-specific nucleases derived from a variety of thermostable organisms are provided. These structure-specific nucleases are used to cleave target-dependent cleavage structures, thereby indicating the presence of specific nucleic acid sequences or specific variations thereof. Disclosed are methods for the detection and characterization of nucleic acid sequences and their variants by using structure-specific 5'-nucleases derived from thermostable DNA polymerases, or the FEN-1, RAD2, or XPG class of nucleases. The enzyme cleaves the target nucleic acid sequence at a structure formed via annealing with 2 pilot oligonucleotide sequences. Also disclosed are methods and devices for the sepn. of nucleic acid mols. based on charge. Also disclosed are methods for the detection of non-target cleavage products via the formation of a complete and activated protein binding region. Isolation of genes for endonuclease FEN-1 from Pyrococcus woesei, Methanococcus jannaschii, Archaeoglobus fulgidus, Methanobacterium thermoautotrophicum, and Pyrococcus furiosus are described. Prepn. of 5'-nucleases by deleting the C-terminal polymerase domain or by point mutations of Taq DNA polymerase, and the prepn. of chimeric enzymes of the FEN-1 endonucleases are also shown. The cleavage method was used for the identification of hepatitis C virus and human ras gene.

REFERENCE COUNT:

7 THERE ARE 7 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L12 ANSWER 19 OF 32 CAPLUS COPYRIGHT 2003 ACS
 ACCESSION NUMBER: 1998:785600 CAPLUS
 DOCUMENT NUMBER: 130:33958
 TITLE: Cleavage of nucleic acid acid using thermostable Methanococcus jannaschii FEN-1 endonucleases
 INVENTOR(S): Kaiser, Michael W.; Lyamichev, Victor I.; Lyamichev, Natasha
 PATENT ASSIGNEE(S): Third Wave Technologies, Inc., USA
 SOURCE: U.S., 330 pp., Cont.-in-part of U.S. Ser. No. 599,491.
 DOCUMENT TYPE: Patent
 LANGUAGE: English
 FAMILY ACC. NUM. COUNT: 12
 PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 5843669	A	19981201	US 1996-757653	19961129
US 5846717	A	19981208	US 1996-599491	19960124
WO 9823774	A1	19980604	WO 1997-US21783	19971126
W: AU, CA, JP, US, US				
RW: AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE				
AU 9855898	A1	19980622	AU 1998-55898	19971126
AU 737449	B2	20010823		
EP 966542	A1	19991229	EP 1997-952237	19971126
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, FI				
JP 2001526526	T2	20011218	JP 1998-524043	19971126
PRIORITY APPLN. INFO.:				
US 1996-599491 A2 19960124				
US 1996-757653 A2 19961129				
US 1996-758314 A2 19961202				
WO 1997-US21783 W 19971126				

AB The present invention relates to means for cleaving a nucleic acid cleavage structure in a site-specific manner. Structure-specific nucleases, including 5' nucleases, thermostable FEN-1 endonucleases and 3' exonucleases, are used to detect and identify target nucleic acids. Methods are provided which allow for the detection specific nucleic acid sequences; these methods permit the detection and identification of mutant and wild-type forms of genes (e.g., human genes) as well as

permit the detection and identification of bacterial and viral pathogens in a sample.

REFERENCE COUNT: 34 THERE ARE 34 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L12 ANSWER 20 OF 32 SCISEARCH COPYRIGHT 2003 ISI (R)
ACCESSION NUMBER: 1998:607277 SCISEARCH

THE GENUINE ARTICLE: 106WC

TITLE:

Engineering an Mg²⁺ site to replace a structurally conserved arginine in the catalytic center of histidyl-tRNA synthetase by computer experiments

AUTHOR:

Arnez J G; Flanagan K; Moras D; Simonson T (Reprint)
UNIVERSITE LOUIS PASTEUR, CNRS, INSERM, INST GENET & BIOL MOL & CELLULAIRE, STRUCT BIOL LAB, BP 163, F-67404

CORPORATE SOURCE:

STRASBOURG ILLKIR, FRANCE (Reprint); UNIVERSITE LOUIS PASTEUR, CNRS, INSERM, INST GENET & BIOL MOL & CELLULAIRE, STRUCT BIOL LAB, F-67404 STRASBOURG ILLKIR, FRANCE

COUNTRY OF AUTHOR:

SOURCE:

PROTEINS-STRUCTURE FUNCTION AND GENETICS, (15 AUG 1998)
Vol. 32, No. 3, pp. 362-380.

Publisher: WILEY-LISS, DIV JOHN WILEY & SONS INC, 605 THIRD AVE, NEW YORK, NY 10158-0012.

ISSN: 0887-3585.

DOCUMENT TYPE:

Article; Journal

FILE SEGMENT:

LIFE

LANGUAGE:

English

REFERENCE COUNT:

45

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS

AB Histidyl-tRNA synthetase (HisRS) differs from other class II aminoacyl-tRNA synthetases (aaRS) in that it harbors an arginine at a position where the others bind a catalytic Mg²⁺ ion. In computer experiments, four mutants of HisRS from *Escherichia coli* were engineered by removing the arginine and introducing a Mg²⁺ ion and residues from seryl-tRNA synthetase (SerRS) that are involved in Mg²⁺ binding. The mutants recreate an active site carboxylate pair conserved in other class II aaRSs, in two possible orders: Glu-Asp or Asp-Glu, replacing Glu-Thr in native HisRS. The mutants were simulated by molecular dynamics in complex with histidyl-adenylate. As controls, the native HisRS was simulated in complexes with histidine, histidyl-adenylate, and histidinol. The native structures sampled were in good agreement with experimental structures and biochemical data. The two mutants with the Glu-Asp sequence showed significant differences in active site structure and Mg²⁺ coordination from SerRS. The others were more similar to SerRS, and one of them was analyzed further through simulations in complex with histidine, and His+ATP. The latter complex sampled two Mg²⁺ positions, depending on the conformation of a loop anchoring the second carboxylate. The lowest energy conformation led to an active site geometry very similar to SerRS, with the principal Mg²⁺ bridging the alpha- and beta-phosphates, the first carboxylate (Asp) coordinating the ion through a water molecule, and the second (Glu) coordinating it directly. This mutant is expected to be catalytically active and suggests a basis for the previously unexplained conservation of the active site Asp-Glu pair in class II aaRSs other than HisRS. Proteins 32:362-380, 1998. (C) 1998 Wiley-Liss, Inc.

L12 ANSWER 21 OF 32 CAPLUS COPYRIGHT 2003 ACS
ACCESSION NUMBER: 1997:260104 CAPLUS

DOCUMENT NUMBER:

126:260880

TITLE:

5' nucleases derived from thermostable DNA polymerases and their use in a nucleic acid detection method

INVENTOR(S):

Dahlberg, James E.; Lyamichev, Victor I.; Brow, Mary Ann D.

PATENT ASSIGNEE(S):

Third Wave Technologies, Inc., USA

SOURCE:

U.S., 93 pp., Cont.-in-part of U.S. 5,541,311.

DOCUMENT TYPE:

CODEN: USXXAM

LANGUAGE:

Patent

FAMILY ACC. NUM. COUNT:

English

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 5614402	A	19970325	US 1994-254359	19940606
US 5422253	A	19950606	US 1992-986330	19921207
US 5541311	A	19960730	US 1993-73384	19930604
US 5837450	A	19981117	US 1995-471066	19950606
US 5843654	A	19981201	US 1995-484956	19950607
US 5719028	A	19980217	US 1997-789079	19970206
US 5888780	A	19990330	US 1997-802233	19970219
US 2003054338	A1	20030320	US 2001-940925	20010828
PRIORITY APPLN. INFO.:				
			US 1992-986330	A2 19921207
			US 1993-73384	A2 19930604
			US 1994-254359	A3 19940606
			US 1994-337164	B2 19941109
			US 1995-402601	A2 19950309
			US 1995-484956	A2 19950607
			US 1995-520946	A1 19950830
			US 1997-789079	A2 19970206
			US 1997-802233	A2 19970219
			US 2000-655378	A3 20000905

AB Derivs. of thermostable **DNA polymerases** that retain their 5'-nuclease activity but lack polymerase are described for use in a nucleic acid detection system. The nuclease activity cleaves the single-stranded moiety of a Y-shaped structure and so is of use in selected cleavage of reporter sequences in a hybridization assay that includes two 5'-nuclease-dependent cleavage and amplification steps. The presence of the target sequence is demonstrated by the release of the reporter moiety from sequences immobilized on a carrier. The ability of the nuclease activity to cleave such structures was shown by the inability of intact Taq polymerase to amplify a hairpin sequence, although the nuclease-free Stoffel fragment could amplify the target sequence. The prepn. and characterization of a no. of polymerase **mutants** for use in these assays is demonstrated. Specific alterations of the *Thermus aquaticus* Taq gene were: a deletion between nucleotides 1601 and 2502 (the end of the coding region), a 4-nucleotide insertion at position 2043, and deletions between nucleotides 1614 and 1848 and between nucleotides 875 and 1778. Three of these derived 5'-nucleases were designated Cleavase BX, Cleavase BB, and Cleavase BN.